Amendments to the Specification

Please amend the title as follows:

~~REAGENT~~ METHOD FOR DIAGNOSIS OF CROHN'S DISEASE

Page 12, lines 10-19, please rewrite as follows:

PCR for differential display method was carried out with anchored oligo (dT) 29-mer (T primers) and non-specific 5' oligonucleotide 25-mer (P primers) (both from Clontech). Ten kinds of P primers and 9 kinds of T primers were appropriately combined to give 90 combinations of different non-specific PCR primer pairs, and using the primer pairs, cDNA was amplified. That is, cDNA synthesized from 0.01 or 0.0025 μg of the total RNA was mixed with 50 μM dNTP mix (Clontech), 1 μM P primer, 1 μM T primer, and Advantage ~~KlenTaq~~ KLENTAQ® Polymerase mix (50X) (Clontech) in a final volume of 10 μl, and was PCR amplified under the following conditions (Thermal cycler MP, Takara).

Page 13, lines 19-31, please rewrite as follows:

The total RNA was extracted from the lesion and non-lesion of six Crohn's disease descending colon excision specimens (excised in a surgical operation at Social Health Insurance Medical Center (Tokyo, Japan)) using a TRIZOL reagent (Life Technologies, MD) following the manual attached thereto. First-strand cDNA was then synthesized from 3 μg of total RNA using 1.5 μg oligo $(dT)_{15}$ primer, 1 mM dNTP mix, and 300 unit MMLV reverse transcriptase (Clontech, CA) in the final volume of 15 μl at 42°C for 1 hour, followed by incubation at 75°C for 10 min. Using the obtained 1st strand cDNA as a template, semi-quantitative RT-PCR was conducted. As a control, GAPDH mRNA was used. Using the respective primer pairs as shown in Table 1, ~~GeneAmp~~ GENEAMP® reagents and ~~AmpliTaq Gold~~ AMPLITAQ GOLD® DNA polymerase (Perkin-Elmer, Norwalk, CT), PCR amplification was performed (Thermal cycler MP, Takara).